Access DB# _____

SEARCH REQUEST FORM

Scientific and Technical Information Center

Requester's Full Name: _____ Examiner # : _____ Date: _____
Art Unit: _____ Phone Number 30_____ Serial Number: _____
Mail Box and Bldg/Room Location: _____ Results Format Preferred (circle): PAPER DISK E-MAIL

If more than one search is submitted, please prioritize searches in order of need.

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Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched. Include the elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the invention. Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc, if known. Please attach a copy of the cover sheet, pertinent claims, and abstract.

Title of Invention: _____

Inventors (please provide full names): _____

_____

Earliest Priority Filing Date: _____

***For Sequence Searches Only* Please include all pertinent information (parent, child, divisional, or issued patent numbers) along with the appropriate serial number.**

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STAFF USE ONLY	Type of Search	Vendors and cost where applicable
Searcher: Richerzhagen	NA Sequence (#) 1	STN
Searcher Phone #: 308-4292	AA Sequence (#)	Dialog
Searcher Location: CM1-6A03	Structure (#)	Questel/Orbit
Date Searcher Picked Up:	Bibliographic	Dr.Link
Date Completed: 4/18	Litigation	Lexis/Nexis
Searcher Prep & Review Time: 19	Fulltext	Sequence Systems Commyn
Clerical Prep Time:	Patent Family	WWW/Internet
Online Time: 8	Other	Other (specify)

PTO-1590 (8-01)

91788

Schreiber, David

From: Steadman, David (AU1652)
Sent: Monday, March 31, 2003 8:20 AM
To: Schreiber, David
Subject: 09/541,462 sequence search request

NAME: David Steadman
AU: 1652
Date:03/31/03
Office: 10D-04
Mailbox: 10D-01

Mr. Schreiber, please search the following sequences in commercial and interference databases:

1) Standard search of SEQ ID NO:1 (polynucleotide) against **nucleic acid** databases.

2) Standard search of SEQ ID NO:2 (polypeptide) against **nucleic acid** databases.

3) Oligo search of SEQ ID NO:1 (polynucleotide) against **nucleic acid** databases.

4) Oligo search of SEQ ID NO:2 (polypeptide) against **nucleic acid** databases.

Please save results to diskette.

Thank you very much.

David J. Steadman
Art Unit 1652
Crystal Mall 1 Room 10D-04
703-308-3934

Pending Nucleic Acid and/or Pending Amino Acid database searches now generate two sets of results. These databases were split into to two parts to reduce the time needed to update the databases daily. The split freed up more machine time for processing searches.

Searches run against the Nucleic Acid Pending database produce two sets of results, with the extensions, **.rnpm** and **.rnpn**
Searches run against the Amino Acid Pending database produce two sets of results, with the extensions, **.rapm** and **.rapn**

The Pending database search results should not be left in the case because they contain data that is confidential.